

Valerio Iebba

List of Publications by Year in descending order

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Version: 2024-02-01

63
papers

4,055
citations

159358

30
h-index

149479

56
g-index

68
all docs

68
docs citations

68
times ranked

5388
citing authors

#	ARTICLE	IF	CITATIONS
1	Intestinal Akkermansia muciniphila predicts clinical response to PD-1 blockade in patients with advanced non-small-cell lung cancer. <i>Nature Medicine</i> , 2022, 28, 315-324.	15.2	225
2	Cancer Induces a Stress Ileopathy Depending on β^2 -Adrenergic Receptors and Promoting Dysbiosis that Contributes to Carcinogenesis. <i>Cancer Discovery</i> , 2022, 12, 1128-1151.	7.7	44
3	Short-chain fatty acids promote the effect of environmental signals on the gut microbiome and metabolome in mice. <i>Communications Biology</i> , 2022, 5, .	2.0	16
4	Microbiota tryptophan metabolism induces aryl hydrocarbon receptor activation and improves alcohol-induced liver injury. <i>Gut</i> , 2021, 70, 1299-1308.	6.1	92
5	Ileal immune tonus is a prognosis marker of proximal colon cancer in mice and patients. <i>Cell Death and Differentiation</i> , 2021, 28, 1532-1547.	5.0	11
6	Ketogenic diet and ketone bodies enhance the anticancer effects of PD-1 blockade. <i>JCI Insight</i> , 2021, 6, .	2.3	143
7	Intestinal microbiota influences clinical outcome and side effects of early breast cancer treatment. <i>Cell Death and Differentiation</i> , 2021, 28, 2778-2796.	5.0	72
8	Multifaceted modes of action of the anticancer probiotic <i>Enterococcus hirae</i> . <i>Cell Death and Differentiation</i> , 2021, 28, 2276-2295.	5.0	18
9	Profiling of Oral Microbiota and Cytokines in COVID-19 Patients. <i>Frontiers in Microbiology</i> , 2021, 12, 671813.	1.5	50
10	Gut microbiota signatures are associated with toxicity to combined CTLA-4 and PD-1 blockade. <i>Nature Medicine</i> , 2021, 27, 1432-1441.	15.2	216
11	<i>Lactobacillus iners</i> and <i>gasseri</i> , <i>Prevotella bivia</i> and HPV Belong to the Microbiological Signature Negatively Affecting Human Reproduction. <i>Microorganisms</i> , 2021, 9, 39.	1.6	22
12	High abundance of genus <i>Prevotella</i> is associated with dysregulation of IFN-I and T cell response in HIV-1-infected patients. <i>Aids</i> , 2020, 34, 1467-1473.	1.0	14
13	Structural Variations of Vaginal and Endometrial Microbiota: Hints on Female Infertility. <i>Frontiers in Cellular and Infection Microbiology</i> , 2020, 10, 350.	1.8	67
14	Cross-reactivity between tumor MHC class II-restricted antigens and an enterococcal bacteriophage. <i>Science</i> , 2020, 369, 936-942.	6.0	217
15	Gut Bacteria Composition Drives Primary Resistance to Cancer Immunotherapy in Renal Cell Carcinoma Patients. <i>European Urology</i> , 2020, 78, 195-206.	0.9	192
16	Chemotherapy-induced ileal crypt apoptosis and the ileal microbiome shape immunosurveillance and prognosis of proximal colon cancer. <i>Nature Medicine</i> , 2020, 26, 919-931.	15.2	118
17	Physiologic colonic uptake of ¹⁸ F-FDG on PET/CT predicts immunotherapy response and gut microbiome diversity in patients with advanced non-small cell lung cancer (NSCLC).. <i>Journal of Clinical Oncology</i> , 2020, 38, 9600-9600.	0.8	0
18	Nasal microbiota in RSV microbiota. , 2019, , .		2

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19	Fecal Microbial Transplantation impact on gut microbiota composition and metabolome, microbial translocation and T-lymphocyte immune activation in recurrent <i>Clostridium difficile</i> infection patients. <i>New Microbiologica</i> , 2019, 42, 221-224.	0.1	7
20	Swimming and rafting of <i>E.coli</i> microcolonies at air-liquid interfaces. <i>MicrobiologyOpen</i> , 2018, 7, e00532.	1.2	7
21	Combining amplicon sequencing and metabolomics in cirrhotic patients highlights distinctive microbiota features involved in bacterial translocation, systemic inflammation and hepatic encephalopathy. <i>Scientific Reports</i> , 2018, 8, 8210.	1.6	63
22	1014 - Microbiota Composition, Metabolic Profiles and Inflammatory Host Response after Fecal Microbiota Transplantation (FMT) for Recurrent <i>Clostridium Difficile</i> Infection. Does Proteobacteria Abundance Predict the Response to FMT?. <i>Gastroenterology</i> , 2018, 154, S-192.	0.6	0
23	Rebuilding the Gut Microbiota Ecosystem. <i>International Journal of Environmental Research and Public Health</i> , 2018, 15, 1679.	1.2	231
24	Gut microbiome composition to predict resistance in renal cell carcinoma (RCC) patients on nivolumab. <i>Journal of Clinical Oncology</i> , 2018, 36, 4519-4519.	0.8	4
25	Behaviour of <i>Bdellovibrio bacteriovorus</i> in the presence of Gram-positive <i>Staphylococcus aureus</i> . <i>New Microbiologica</i> , 2018, 41, 145-152.	0.1	16
26	Optimizing an array of self adapted temperature modulated metal oxide sensors for biomedical application. , 2017, , .		1
27	Exposure of <i>E. coli</i> to DNA-Methylating Agents Impairs Biofilm Formation and Invasion of Eukaryotic Cells via Down Regulation of the N-Acetylneuraminase Lyase <i>NanA</i> . <i>Frontiers in Microbiology</i> , 2016, 7, 147.	1.5	13
28	Mo1927 Mucosa-Associated Microbiota and Promoter Methylation Status of Genes Involved in Immune Response in Crohn's Disease Patients. <i>Gastroenterology</i> , 2016, 150, S818.	0.6	0
29	Uncovering oral <i>Neisseria tropism</i> and persistence using metagenomic sequencing. <i>Nature Microbiology</i> , 2016, 1, 16070.	5.9	68
30	Protective Role of Postbiotic Mediators Secreted by <i>Lactobacillus rhamnosus</i> GG Versus Lipopolysaccharide-induced Damage in Human Colonic Smooth Muscle Cells. <i>Journal of Clinical Gastroenterology</i> , 2016, 50, S140-S144.	1.1	38
31	Bacterial Biofilm in Salivary Gland Stones. <i>Otolaryngology - Head and Neck Surgery</i> , 2016, 154, 449-453.	1.1	17
32	What is new about diet in hepatic encephalopathy. <i>Metabolic Brain Disease</i> , 2016, 31, 1289-1294.	1.4	30
33	Gut microbiota related to <i>Giardia duodenalis</i> , <i>Entamoeba</i> spp. and <i>Blastocystis hominis</i> infections in humans from Cote d'Ivoire. <i>Journal of Infection in Developing Countries</i> , 2016, 10, 1035-1041.	0.5	89
34	Eubiosis and dysbiosis: the two sides of the microbiota. <i>New Microbiologica</i> , 2016, 39, 1-12.	0.1	109
35	OP03 THE PROTECTIVE ROLE OF <i>LACTOBACILLUS RHAMNOSUS</i> GG-DERIVED FACTORS AGAINST LPS-INDUCED DAMAGE OF HUMAN COLONIC SMOOTH MUSCLE CELLS. <i>Journal of Pediatric Gastroenterology and Nutrition</i> , 2015, 61, 509-510.	0.9	1
36	Su1861 Postbiotic Protective Activity of <i>Lactobacillus Rhamnosus</i> GG-Derived Factors on Pathogen Lipopolysaccharide (LPS)-Induced Damage of Human Colonic Smooth Muscle. <i>Gastroenterology</i> , 2015, 148, S-536.	0.6	0

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37	Outbreak of <i>Achromobacter xylosoxidans</i> in an Italian Cystic fibrosis center: genome variability, biofilm production, antibiotic resistance, and motility in isolated strains. <i>Frontiers in Microbiology</i> , 2014, 5, 138.	1.5	46
38	<i>Bdellovibrio bacteriovorus</i> directly attacks <i>Pseudomonas aeruginosa</i> and <i>Staphylococcus aureus</i> Cystic fibrosis isolates. <i>Frontiers in Microbiology</i> , 2014, 5, 280.	1.5	74
39	Microbiota and the gut-liver axis: Bacterial translocation, inflammation and infection in cirrhosis. <i>World Journal of Gastroenterology</i> , 2014, 20, 16795.	1.4	187
40	<i>Escherichia coli</i> Population-Based Study in Pediatric Crohn&€™s Disease. <i>Advances in Microbiology</i> , 2014, 04, 886-889.	0.3	1
41	Cystic Fibrosis Transmembrane Conductance Regulator (CFTR) Allelic Variants Relate to Shifts in Faecal Microbiota of Cystic Fibrosis Patients. <i>PLoS ONE</i> , 2013, 8, e61176.	1.1	83
42	Higher Prevalence and Abundance of <i>Bdellovibrio bacteriovorus</i> in the Human Gut of Healthy Subjects. <i>PLoS ONE</i> , 2013, 8, e61608.	1.1	93
43	Microevolution in <i>fimH</i> Gene of Mucosa-Associated <i>Escherichia coli</i> Strains Isolated from Pediatric Patients with Inflammatory Bowel Disease. <i>Infection and Immunity</i> , 2012, 80, 1408-1417.	1.0	49
44	Gut Microbiota and the Immune System: An Intimate Partnership in Health and Disease. <i>International Journal of Immunopathology and Pharmacology</i> , 2012, 25, 823-833.	1.0	30
45	Plasmid-mediated fluoroquinolone resistance determinants in <i>Escherichia coli</i> from community uncomplicated urinary tract infection in an area of high prevalence of quinolone resistance. <i>European Journal of Clinical Microbiology and Infectious Diseases</i> , 2012, 31, 1917-1921.	1.3	21
46	A potential role of <i>Escherichia coli</i> pathobionts in the pathogenesis of pediatric inflammatory bowel disease. <i>Canadian Journal of Microbiology</i> , 2012, 58, 426-432.	0.8	29
47	Assessment Of Daily Physical Activity In Patients With Cystic Fibrosis. , 2012, , .		0
48	Influence of oral conditions on colonization by highly toxigenic <i>Staphylococcus aureus</i> strains. <i>Oral Diseases</i> , 2012, 18, 402-409.	1.5	20
49	Diffusion of meticillin-resistant <i>Staphylococcus aureus</i> USA300 strains in central Italy. <i>International Journal of Antimicrobial Agents</i> , 2011, 37, 339-346.	1.1	9
50	Swimming with an Image. <i>Physical Review Letters</i> , 2011, 106, 038101.	2.9	217
51	Gut Microbiota and Pediatric Disease. <i>Digestive Diseases</i> , 2011, 29, 531-539.	0.8	34
52	Microbiological and molecular characterization of nosocomial and community <i>Staphylococcus aureus</i> isolates. <i>Epidemiology and Infection</i> , 2011, 139, 613-622.	1.0	9
53	JC Viral Reactivation in a Pediatric Patient with Crohn's Disease. <i>International Journal of Immunopathology and Pharmacology</i> , 2010, 23, 955-959.	1.0	8
54	A distinctive 'microbial signature' in celiac pediatric patients. <i>BMC Microbiology</i> , 2010, 10, 175.	1.3	201

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55	<i>Achromobacter xylosoxidans</i> Genomic Characterization and Correlation of Randomly Amplified Polymorphic DNA Profiles of Cystic Fibrosis Patients. <i>Journal of Clinical Microbiology</i> , 2010, 48, 1035-1039.	1.8	14
56	Bacterial ratchet motors. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 9541-9545.	3.3	559
57	T1811 Dominant Intestinal Mucosa-Associated Microbiota in Pediatric Patients With Inflammatory Bowel Disease. <i>Gastroenterology</i> , 2010, 138, S-583-S-584.	0.6	0
58	Dominant genotypes in mucosa-associated <i>Escherichia coli</i> strains from pediatric patients with inflammatory bowel disease. <i>Inflammatory Bowel Diseases</i> , 2009, 15, 661-672.	0.9	38
59	Effect of lactoferricin on fluoroquinolone susceptibility of uropathogenic <i>Escherichia coli</i> . <i>Journal of Antibiotics</i> , 2009, 62, 109-111.	1.0	5
60	The Microbiota in Inflammatory Bowel Disease in Different Age Groups. <i>Digestive Diseases</i> , 2009, 27, 252-258.	0.8	56
61	T1274 Dominant Mucosa-Associated Microbiota in Celiac Children At Diagnosis and After GFD. <i>Gastroenterology</i> , 2008, 134, A-520.	0.6	0
62	W1188 Dominant Genotypes in Mucosa-Associated <i>Escherichia coli</i> Strains from Pediatric Patients with Inflammatory Bowel Disease. <i>Gastroenterology</i> , 2008, 134, A-651.	0.6	0
63	Virulence Traits in <i>Escherichia Coli</i> Strains Isolated from Outpatients with Urinary Tract Infections. <i>International Journal of Immunopathology and Pharmacology</i> , 2008, 21, 715-723.	1.0	2